

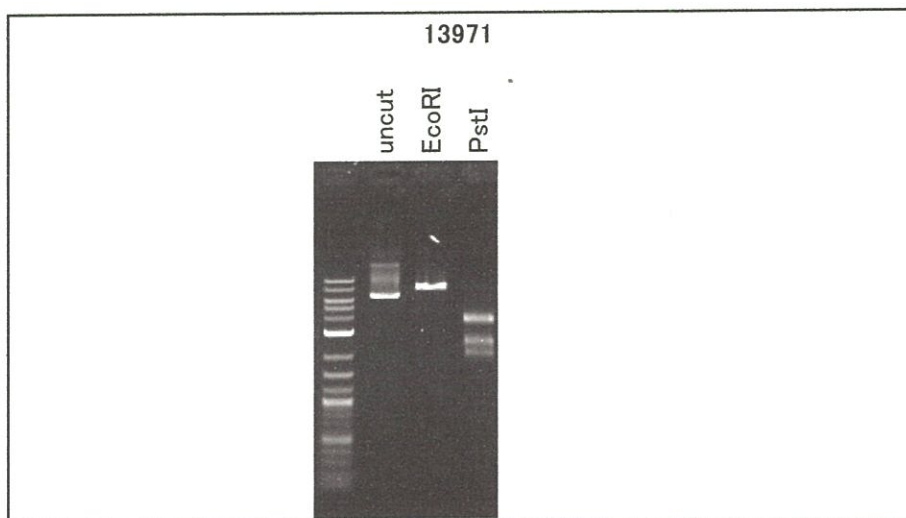
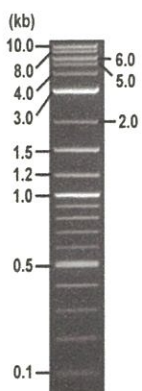


**RIKEN DNA BANK**

clone name : pCIneoLuc-PP1A

- Clone ID : RDB \_ 13971
- Lot : 13971 \_ A6Bt
- DNA Concentration : 25 nanogram/microliter
- Volume : 40 microliter
- Form : DNA solution in TE buffer
- Host : DH5 alpha
- Culture : LB medium
- Antibiotics : 100 microgram/ml Ampicillin
- Purification : QIAGEN QIAprep Spin Miniprep kit
- Digestion by restriction enzyme

2-Log DNA Ladder  
(NEB#N3200L),  
125 ng/well



Electrophoresis : 88 nanogram DNA per lane ; 1% agarose gel , 1 x TAE Buffer

Restriction enzyme	Expected size of fragment
<u>EcoRI</u>	<u>8.1</u> kbp
<u>PstI</u>	<u>3.7, 2.5, 2.0</u> kbp
_____	_____ kbp
_____	_____ kbp
_____	_____ kbp

● Confirmation of the insertion sequence

Sequence name	Primer name	Sequence name	Primer name
Sequence - A	CMV-Forward	Sequence - E	-
Sequence - B	Luc-C_F	Sequence - F	-
Sequence - C	pAxCALNL_F1	Sequence - G	-
Sequence - D	SV40pro_F_V2	Sequence - H	-

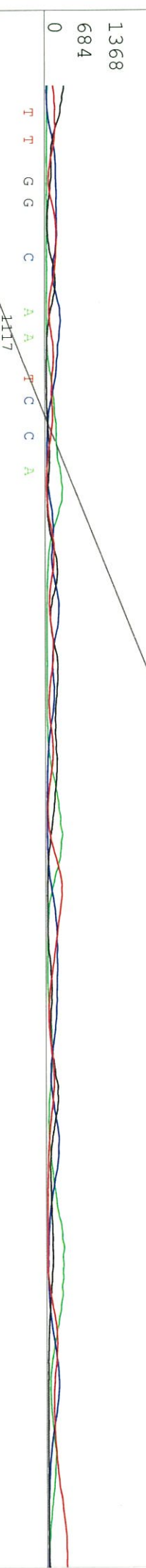
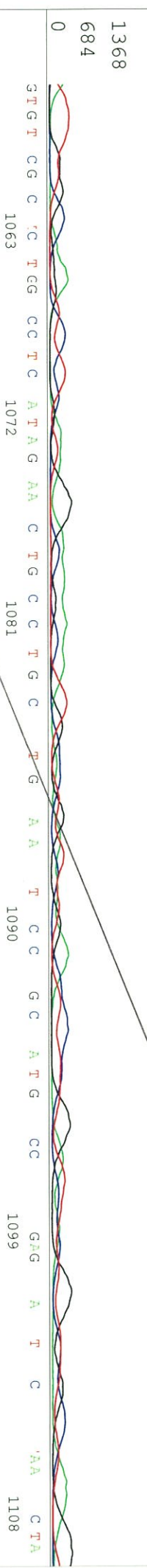
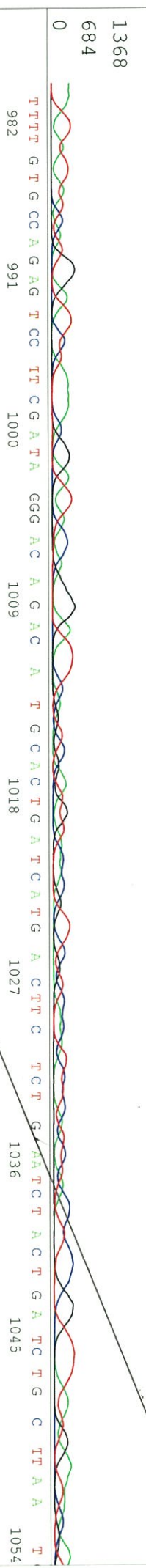
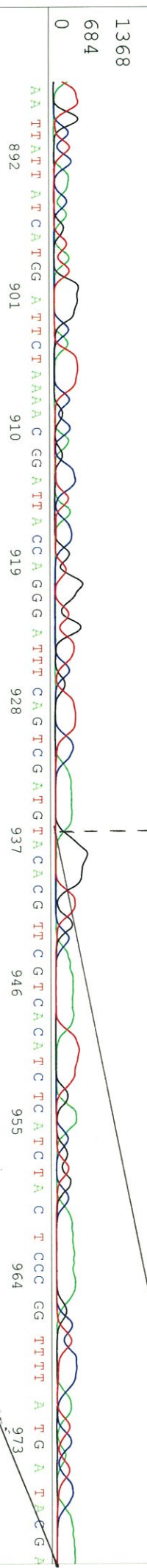
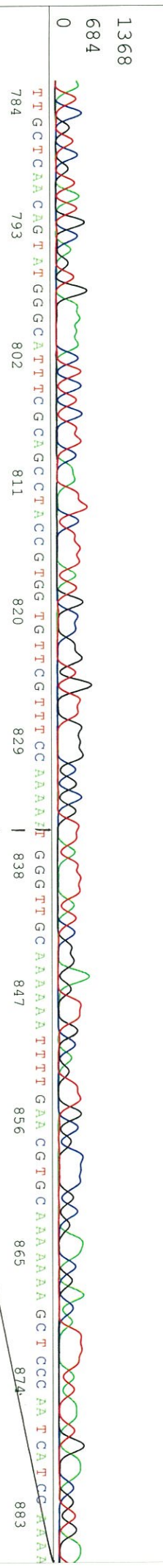


APPROVED BY :





ATCGTCGTTATGCAAGTGAAGAAACCTCTCTCAATTCCTTTATGCGCCGGTGTGGGCGCGTTATTATTCCGGAGTTGCAAGTTGCCCGCCGAAAGCAATTTATAATGAACGGTGA



Printed on: Wed Mar 09, 2016 01:10PM, JST  
Electropherogram Data Page 2 of 2



















S/N G:383 A:342 T:290 C:432

primer name  $\varnothing$  : SV40pro\_F\_V2

KB\_3500\_POP7\_BDTV3.mob

Mar 09, 2016 10:45AM, JST

KB.bcp

5'-TCGGCCTCTGAGCTATTC-3'

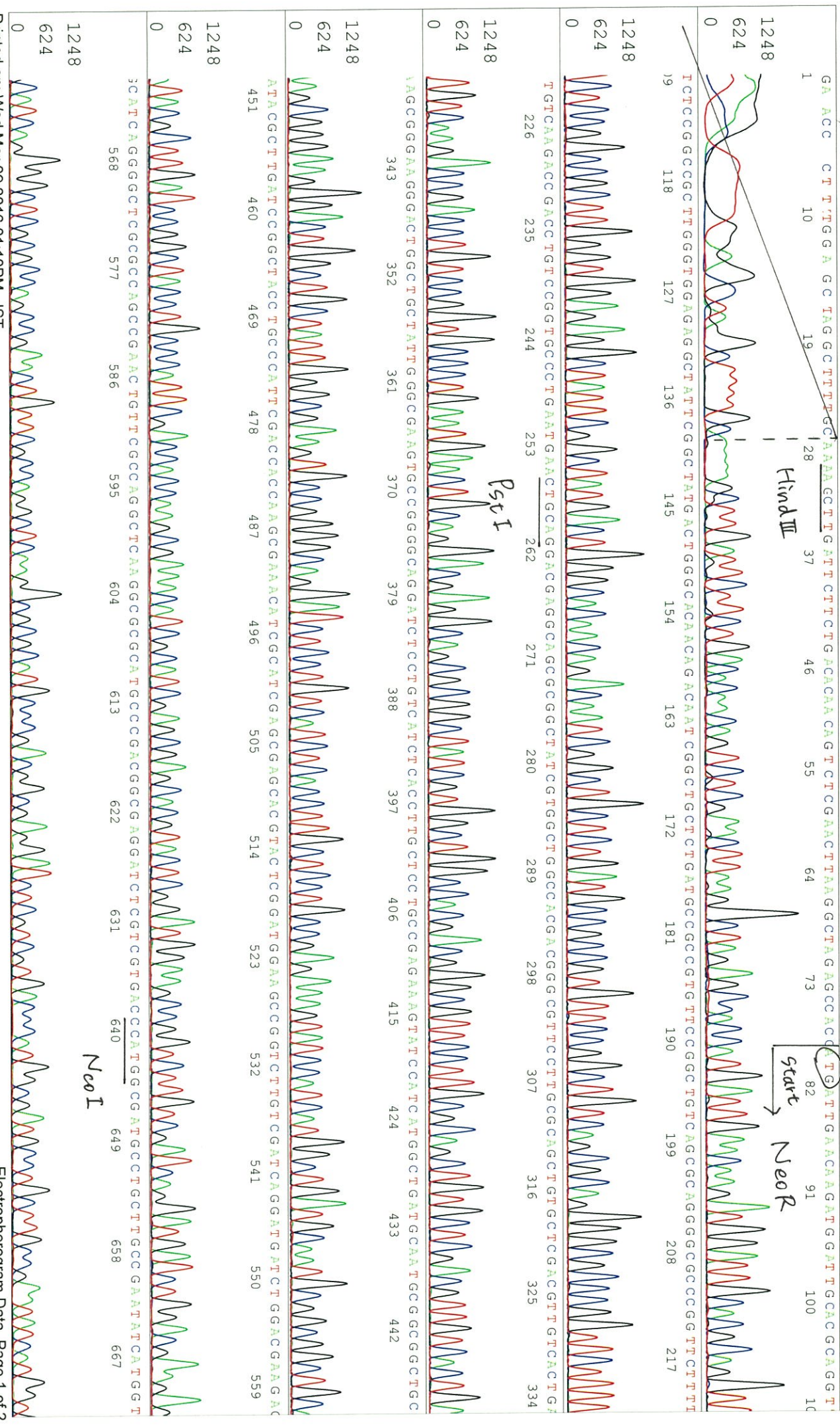
Pts 1401 to 12972 Pk1 Loc:1378

Mar 09, 2016 11:11AM, JST

KB 1.4:1.8 Cap:19

Version 6.0 HISQV Bases: 984

Plate Name: 20160309\_GNP





GGAAAATGGCCGCTTTTCTGATTCATCGACTGTGGCCGGCTGGGTGGCCGGAACCCCTTATCAGGACATAGCCGTTGGCTACCCTGATATTGCTGAGAGAGCTTG GCG



3CGAATGGGCTGACCGCTTCTCGTGTGCTTTACGGTATCGCCGCTCCCGA TTCCGACCGCATCCGCTTCTGATAGGAGTTCTTCTGAGCGGGACTC

NeoR →

TGGGGGTTCGAATGACCGAACGAGCGCCAA CCGCCATCAGGATGGCCGCAATAAATAATCTTTATTCTCA TTACATCTGGTG TGGG TTT

TTTGTGTGAA TCGAATA GCGATCCGGCTATGG TGCATCTCAAG TACC ATCTGCCTTCTG ATGCCCGCA TAG TT AAA G C

AG CCCC GG ACA C C (A C AA C A C C G C TG A C G C C TTG ACC A C C G C C T GA TCC TG C T C C G G C

AA T T C C G C G C T T